STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/537,449A
Source: IFW16
Date Processed by STIC: 10/5/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):** U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 10/537,449A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4	Non-ASCII	The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)
11	Use of <220>	Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n/Xaa	"n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid

IFW16

RAW SEQUENCE LISTING DATE: 10/05/2006
PATENT APPLICATION: **US/10/537,449A** TIME: 11:38:44

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\10052006\J537449A.raw**

```
      3 <110> APPLICANT: Technische Universitat Dresden
      5 <120> TITLE OF INVENTION: Polynucleotides Targeted Against Htert and Use Thereof
      7 <130> FILE REFERENCE: 101215-189-2
      9 <140> CURRENT APPLICATION NUMBER: 10/537,449A
C--> 10 <141> CURRENT FILING DATE: 2006-01-09
     12 <160> NUMBER OF SEQ ID NOS: 18
     14 <170> SOFTWARE: PatentIn Ver. 2.1
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 75
     18 <212> TYPE: DNA
     19 <213> ORGANISM: Homo sapiens
     21 <220> FEATURE:
     22 <221> NAME/KEY: mRNA
     23 <222> LOCATION: (1)..(75)
     24 <223> OTHER INFORMATION: subunit 2176-2250 of hTERT
W--> 26 <307> DATE:
     27 <308> DATABASE ACCESSION NO: AF015950
W--> 28 <309> DATABASE ENTRY DATE: AUG 1997
     30 <400> SEQUENCE: 1
     31 ctttgtcaag gtggatgtga cgggcgcgta cgacaccatc ccccaggaca ggctcacgga 60
     32 ggtcatcgcc agcat                                                  75
     35 <210> SEQ ID NO: 2
     36 <211> LENGTH: 98
     37 <212> TYPE: DNA
     38 <213> ORGANISM: Homo sapiens
     40 <220> FEATURE:
     41 <221> NAME/KEY: mRNA
     42 <222> LOCATION: (1)..(98)
     43 <223> OTHER INFORMATION: subunit 2296-2393 of hTERT
W--> 45 <307> DATE:
     46 <308> DATABASE ACCESSION NO: AF015950
W--> 47 <309> DATABASE ENTRY DATE: AUG 1997
     49 <400> SEQUENCE: 2
     50 ccagaaggcc gcccatgggc acgtccgcaa ggccttcaag agccacgtct ctaccttgac 60
     51 agacctccag ccgtacatgc gacagttcgt ggctcacc                         98
     54 <210> SEQ ID NO: 3
     55 <211> LENGTH: 23
     56 <212> TYPE: DNA
     57 <213> ORGANISM: Homo sapiens
     59 <220> FEATURE:
     60 <221> NAME/KEY: mRNA
     61 <222> LOCATION: (1)..(23)
     62 <223> OTHER INFORMATION: subunit 2183-2205 of hTERT
```

Does Not Comply
Corrected Diskette Needed

insert <3007

delete

add hyphen AUG-1997 (global error)

insert <3007

insert this mandatory numeric identifier above publication information. <3007 is a "header" only. It never has a response.

insert ↓ <3007

RAW SEQUENCE LISTING DATE: 10/05/2006
PATENT APPLICATION: US/10/537,449A TIME: 11:38:44

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\10052006\J537449A.raw

W--> 64 ~~<307> DATE:~~
65 <308> DATABASE ACCESSION NO: ~~AF015950~~
W--> 66 **<309> DATABASE ENTRY DATE: AUG 1997**
68 <400> SEQUENCE: 3
69 aaggtggatg tgacgggcgc gta 23
72 <210> SEQ ID NO: 4
73 <211> LENGTH: 20
74 <212> TYPE: DNA
75 <213> ORGANISM: Homo sapiens
77 <220> FEATURE:
78 <221> NAME/KEY: mRNA
79 <222> LOCATION: (1)..(20)
80 <223> OTHER INFORMATION: subunit 2206-2225 of hTERT <3007
W--> 82 ~~<307> DATE:~~
83 <308> DATABASE ACCESSION NO: AF015950
W--> 84 **<309> DATABASE ENTRY DATE: AUG 1997**
86 <400> SEQUENCE: 4
87 cgacaccatc ccccaggaca 20
90 <210> SEQ ID NO: 5
91 <211> LENGTH: 20
92 <212> TYPE: DNA
93 <213> ORGANISM: Homo sapiens
95 <220> FEATURE:
96 <221> NAME/KEY: mRNA
97 <222> LOCATION: (1)..(20)
98 <223> OTHER INFORMATION: subunit 2315-2334 of hTERT <3007
W--> 100 ~~<307> DATE:~~
101 <308> DATABASE ACCESSION NO: ~~AF015950~~
W--> 102 **<309> DATABASE ENTRY DATE: AUG 1997**
104 <400> SEQUENCE: 5
105 cacgtccgca aggccttcaa 20
108 <210> SEQ ID NO: 6
109 <211> LENGTH: 20
110 <212> TYPE: DNA
111 <213> ORGANISM: Homo sapiens
113 <220> FEATURE:
114 <221> NAME/KEY: mRNA
115 <222> LOCATION: (1)..(20)
116 <223> OTHER INFORMATION: subunit 2317-2336 of hTERT <3007
W--> 118 ~~<307> DATE:~~
119 <308> DATABASE ACCESSION NO: AF015950
W--> 120 **<309> DATABASE ENTRY DATE: AUG 1997**
122 <400> SEQUENCE: 6
123 cgtccgcaag gccttcaaga 20
126 <210> SEQ ID NO: 7
127 <211> LENGTH: 23
128 <212> TYPE: DNA
129 <213> ORGANISM: Homo sapiens
131 <220> FEATURE:

RAW SEQUENCE LISTING DATE: 10/05/2006
PATENT APPLICATION: US/10/537,449A TIME: 11:38:44

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\10052006\J537449A.raw

132 <221> NAME/KEY: mRNA
133 <222> LOCATION: (1)..(23)
134 <223> OTHER INFORMATION: subunit 2324-2346 of hTERT <2207
W--> 136 ~~<307> DATE:~~
137 <308> DATABASE ACCESSION NO: AF015950
W--> 138 **<309> DATABASE ENTRY DATE: AUG 1997**
140 <400> SEQUENCE: 7
141 aaggccttca agagccacgt ctc 23
144 <210> SEQ ID NO: 8
145 <211> LENGTH: 20
146 <212> TYPE: DNA
147 <213> ORGANISM: Homo sapiens
149 <220> FEATURE:
150 <221> NAME/KEY: mRNA
151 <222> LOCATION: (1)..(20)
152 <223> OTHER INFORMATION: subunit 2331-2350 hTERT <3007
W--> 154 ~~<307> DATE:~~
155 <308> DATABASE ACCESSION NO: AF015950
W--> 156 **<309> DATABASE ENTRY DATE: AUG 1997**
158 <400> SEQUENCE: 8
159 tcaagagcca cgtctctacc 20
162 <210> SEQ ID NO: 9
163 <211> LENGTH: 20
164 <212> TYPE: DNA
165 <213> ORGANISM: Homo sapiens
167 <220> FEATURE:
168 <221> NAME/KEY: mRNA
169 <222> LOCATION: (1)..(20)
170 <223> OTHER INFORMATION: subunit 2333-2352 of hTERT <3007
W--> 172 ~~<307> DATE:~~
173 <308> DATABASE ACCESSION NO: AF015950
W--> 174 **<309> DATABASE ENTRY DATE: AUG 1997**
176 <400> SEQUENCE: 9
177 aagagccacg tctctacctt 20
180 <210> SEQ ID NO: 10
181 <211> LENGTH: 20
182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: hTERT-AS AStel 2206-2225 <3007
W--> 188 ~~<307> DATE:~~
189 <308> DATABASE ACCESSION NO: AF015950
W--> 190 **<309> DATABASE ENTRY DATE: AUG 1997**
192 <400> SEQUENCE: 10
193 tgtcctgggg gatggtgtcg 20
196 <210> SEQ ID NO: 11
197 <211> LENGTH: 20
198 <212> TYPE: DNA
199 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING DATE: 10/05/2006
PATENT APPLICATION: US/10/537,449A TIME: 11:38:44

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\10052006\J537449A.raw

201 <220> FEATURE:
202 <223> OTHER INFORMATION: hTERT-AS AStel 2315-2334 (3007
W--> 204 ~~<307> DATE:~~
205 <308> DATABASE ACCESSION NO: AF015950
W--> 206 **<309> DATABASE ENTRY DATE: AUG 1997**
208 <400> SEQUENCE: 11
209 ttgaaggcct tgcggacgtg 20
212 <210> SEQ ID NO: 12
213 <211> LENGTH: 20
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:
218 <223> OTHER INFORMATION: hTERT-AS AStel 2317-2336 (3007
W--> 220 ~~<307> DATE:~~
221 <308> DATABASE ACCESSION NO: AF015950
W--> 222 **<309> DATABASE ENTRY DATE: AUG 1997**
224 <400> SEQUENCE: 12
225 tcttgaaggc cttgcggacg 20
228 <210> SEQ ID NO: 13
229 <211> LENGTH: 20
230 <212> TYPE: DNA
231 <213> ORGANISM: Artificial Sequence
233 <220> FEATURE:
234 <223> OTHER INFORMATION: hTERT-AS AStel 2331-2350 (3007
W--> 236 ~~<307> DATE:~~
237 <308> DATABASE ACCESSION NO: AF015950
W--> 238 **<309> DATABASE ENTRY DATE: AUG 1997**
240 <400> SEQUENCE: 13
241 ggtagagacg tggctcttga 20
244 <210> SEQ ID NO: 14
245 <211> LENGTH: 20
246 <212> TYPE: DNA
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <223> OTHER INFORMATION: hTERT-AS AStel 2333-2352 (3007
W--> 252 **<307> DATE:**
253 <308> DATABASE ACCESSION NO: AF015950
W--> 254 **<309> DATABASE ENTRY DATE: AUG 1997**
256 <400> SEQUENCE: 14
257 aaggtagaga cgtggctctt 20
260 <210> SEQ ID NO: 15
261 <211> LENGTH: 20
262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION: NS-K2 what is its source? (see item 11 on Error Summary Sheet)
268 <300> PUBLICATION INFORMATION:
269 <308> DATABASE ACCESSION NO: AF015950
W--> 270 **<309> DATABASE ENTRY DATE: AUG 1997**

RAW SEQUENCE LISTING DATE: 10/05/2006
PATENT APPLICATION: US/10/537,449A TIME: 11:38:44

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\10052006\J537449A.raw

272 <400> SEQUENCE: 15
273 cagtctcagt actgaagctg 20
276 <210> SEQ ID NO: 16
277 <211> LENGTH: 20
278 <212> TYPE: DNA
279 <213> ORGANISM: Artificial Sequence
281 <220> FEATURE:
282 <223> OTHER INFORMATION: NS-K3 what is its source?
284 <300> PUBLICATION INFORMATION:
285 <308> DATABASE ACCESSION NO: AF015950
W--> 286 <309> DATABASE ENTRY DATE: AUG 1997
288 <400> SEQUENCE: 16
289 cagcttcagt actgagactg 20
292 <210> SEQ ID NO: 17
293 <211> LENGTH: 501
294 <212> TYPE: DNA
295 <213> ORGANISM: Homo sapiens
297 <220> FEATURE:
298 <221> NAME/KEY: mRNA
299 <222> LOCATION: (1)..(501)
300 <223> OTHER INFORMATION: subunit 2000-2500 of hTERT
302 <300> PUBLICATION INFORMATION:
303 <308> DATABASE ACCESSION NO: AF015950
W--> 304 <309> DATABASE ENTRY DATE: AUG 1997
306 <400> SEQUENCE: 17
307 aagagggccg agcgtctcac ctcgagggtg aaggcactgt tcagcgtgct caactacgag 60
308 cgggcgcggc gccccggcct cctgggcgcc tctgtgctgg gcctggacga tatccacagg 120
309 gcctggcgca ccttcgtgct gcgtgtgcgg gcccaggacc cgccgcctga gctgtacttt 180
310 gtcaaggtgg atgtgacggg cgcgtacgac accatccccc aggacaggct cacggaggtc 240
311 atcgccagca tcatcaaacc ccagaacacg tactgcgtgc gtcggtatgc cgtggtccag 300
312 aaggccgccc atgggcacgt ccgcaaggcc ttcaagagcc acgtctctac cttgacagac 360
313 ctccagccgt acatgcgaca gttcgtggct cacctgcagg agaccagccc gctgagggat 420
314 gccgtcgtca tcgagcagag ctcctccctg aatgaggcca gcagtggcct cttcgacgtc 480
315 ttcctacgct tcatgtgcca c 501
318 <210> SEQ ID NO: 18
319 <211> LENGTH: 4015
320 <212> TYPE: DNA
321 <213> ORGANISM: Homo sapiens
323 <220> FEATURE:
324 <221> NAME/KEY: mRNA
325 <222> LOCATION: (1)..(4015)
326 <223> OTHER INFORMATION: hTERT
328 <300> PUBLICATION INFORMATION:
329 <308> DATABASE ACCESSION NO: AF015950
W--> 330 <309> DATABASE ENTRY DATE: AUG 1997
332 <400> SEQUENCE: 18
333 gcagcgctgc gtcctgctgc gcacgtggga agccctggcc ccggccaccc ccgcgatgcc 60
334 gcgcgctccc cgctgccgag ccgtgcgctc cctgctgcgc agccactacc gcgaggtgct 120
335 gccgctggcc acgttcgtgc ggcgcctggg gccccagggc tggcggctgg tgcagcgcgg 180

VERIFICATION SUMMARY DATE: 10/05/2006
PATENT APPLICATION: **US/10/537,449A** TIME: 11:38:45

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\10052006\J537449A.raw**

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:26 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:1
L:28 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:1
L:45 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:2
L:47 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:2
L:64 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:3
L:66 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:3
L:82 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:4
L:84 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:4
L:100 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:5
L:102 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:5
L:118 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:6
L:120 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:6
L:136 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:7
L:138 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:7
L:154 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:8
L:156 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:8
L:172 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:9
L:174 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:9
L:188 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:10
L:190 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:10
L:204 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:11
L:206 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:11
L:220 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:12
L:222 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:12
L:236 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:13
L:238 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:13
L:252 M:285 W: Invalid Journal Date Format:Use YYYY-MM-DD,Mon-YYYY,Season-YYYY,or YYYY, SEQ:14
L:254 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:14
L:270 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:15
L:286 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:16
L:304 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:17
L:330 M:286 W: Invalid Database Entry: Valid Formats YYYY-MM-DD or Mon-YYYY, SEQ ID:18